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RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/083,246

DATE: 08/08/2002

TIME: 10:10:21

Input Set : A:\EP.txt

Output Set: N:\CRF3\08082002\J083246.raw

3 <110> APPLICANT: Athena Diagnostics

5 <120> TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR GENETIC ANALYSIS OF POLYCYSTIC

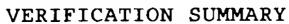
KIDNEY

- DISEASE 6
- 8 <130> FILE REFERENCE: 1133/2002
- 10 <140> CURRENT APPLICATION NUMBER: US 10/083,246
- 11 <141> CURRENT FILING DATE: 2002-02-26
- 13 <160 > NUMBER OF SEQ ID NOS: 168
- 15 <170> SOFTWARE: PatentIn version 3.1

ERRORED SEQUENCES

- 2975 <210> SEQ ID NO: 168
- 2976 <211> LENGTH: (20) Found 22
- 2977 <212> TYPE: DNA
- 2978 <213> ORGANISM: Artificial Sequence
- 2980 <220> FEATURE:
- 2981 <221> NAME/KEY: misc_feature
- 2982 <222> LOCATION: (1)..(22)
- 2983 <223> OTHER INFORMATION: Synthetic primer
- 2987 <400> SEQUENCE: 168

2988 aaatacaact gtcagcaaca ta
E--> 2989 73 - (\text{tht}



PATENT APPLICATION: US/10/083,246

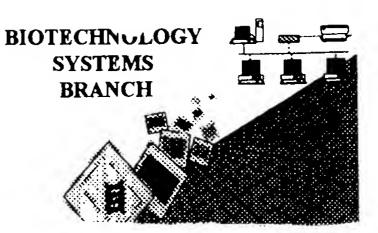
DATE: 08/08/2002 TIME: 10:10:22

Input Set : A:\EP.txt

Output Set: N:\CRF3\08082002\J083246.raw

L:273 M:259 W: Allowed number of lines exceeded, <223> Other Information: L:299 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:660 L:319 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:1260 L:331 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:1620 L:341 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:1920 L:351 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:2220 L:353 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:2280 L:363 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:2580 L:373 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:2880 L:375 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:2940 L:389 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:3360 L:397 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:3600 L:399 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:3660 L:405 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:3840 L:413 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:4080 L:421 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:4320 L:423 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:4380 L:465 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:5640 L:467 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:5700 L:497 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:6600 L:499 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:6660 L:501 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:6720 L:2989 M:254 E: No. of Bases conflict, this line has no nucleotides. L:2989 M:252 E: No. of Seq. differs, <211> LENGTH:Input:20 Found:22 SEQ:168

0500



RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

10/083, 246
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8/8/02

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Revised 01/29/2002